RAW SEQUENCE LISTING

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Application Serial Number: 10/570/047
Source: 1500
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IFW#

RAW SEQUENCE LISTING DATE: 03/08/2006
PATENT APPLICATION: US/10/570,047 TIME: 13:21:03

Input Set : A:\39363a.txt

3 <110> APPLICANT: Jessberger, et al.

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5 <120> TITLE OF INVENTION: METHODS FOR IDENTIFYING, TREATING, AND INDUCING INFERTILITY USING
           SMC1 BETA
   8 <130> FILE REFERENCE: 29636/39363A
-> 10 <140> CURRENT APPLICATION NUMBER: US/10/570,047
-> 10 <141> CURRENT FILING DATE: 2006-02-28
  10 <150> PRIOR APPLICATION NUMBER: US 60/499,317
  11 <151> PRIOR FILING DATE: 2003-08-29
  13 <160> NUMBER OF SEO ID NOS: 13
  15 <170> SOFTWARE: PatentIn version 3.2
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Input Set : A:\39363a.txt

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Output Set: N:\CRF4\03072006\J570047.raw

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RAW SEQUENCE LISTING DATE: 03/08/2006 PATENT APPLICATION: US/10/570,047 TIME: 13:21:03

Input Set : A:\39363a.txt

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312	36-4	17 - L	- 1 -	580	**- 7	- 1 -	T	m1	585	D1	D	~1 ···		590	.	**- 3
	мет	мет		Asp	vaı	тте	гуѕ		GIN	Pne	Pro	GIn		ьуѕ	Lys	vaı
316	т10	C1 n	595	17a]	Crra	C1	7 ~~	600	T 011	77.7	Crra	C1	605	37-7	C1	C1.,
320	116	610	PIIE	vai	Cys	GIY	615	GIY	ьеи	vai	Cys	620	IIIL	vaı	Glu	Giu
	Δla		Hic	Tla	בומ	Dhe		Glv	Pro	Glu	Δra		Luc	Δla	Val	Δ] =
	625	Arg	1113	110	HΙα	630	Gry	Gry	110	GIU	635	nr 9	цуз	nια	vai	640
		Asp	Glv	Thr	Len		Len	Lvs	Ser	Glv		Tle	Ser	Glv	Gly	
328		p			645			_,,	501	650	•		501	017	655	001
	Ser	Asp	Leu	Lvs		Lvs	Ala	Leu	Cvs		asp	Glu	Lvs	Glu	Leu	His
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360	~1	770	C1	7	T7 -	7	775	nh -	~ 7	7	T	780	17c 3	T	~ 1	~1 -
		val	GIU	ASI	тте	_	GIU	rne	GIU	ASN	_	HIS	val	гда	Gln	
	785 Gln	Gl.	7\ α∽	7\ <>~	<u>ما</u> ت	790	7.22	Ten	G1.,	Dho	795	Tara	G1 ~	Tura	Th~	800
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	Glu	Leu	Met	Val	Lys			Gln	Ile	Lys	Glu		Leu	Ala	Thr	Gln
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392	1703	Ile	т1.	900	C1	Cor	T 011	C1	905	T	T 011	Τ 011	~1	910		7 an
396	vai	TTE	915	GIII	Gry	Ser	пеп	920	GIII	цуъ	цец	пец	925	пуъ	птъ	ASII
	Leu	Leu		Asp	Cvs	Lvs	Val		Asp	Ile	Asp	Ile		Leu	Val	Leu
400		930			-1-	-1-	935					940				
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	Asp	Tyr	Ser		Leu	Arg	Glu	Asp		Lys	Ala	Leu	Gln		_	Lys
412	~1	17 7	<i>α</i> 1	980	TT	т	mh	T	985	. т		. (1)	17-	990		an (1) n
415	GIU	vaı	995	Ala	HIS	ьeu	THE	1000		а це	ı Gı	1 G1:	n va 10		ıa s	er Gln
			993					1000	,				10	0.5		
419	Clu	Acn	Thr	T.=1	ı T.e.ı	1 T.370	Thr	~ т·}	nar Δ΄	la Di	ro A	en L	۵11	Δτα	Δla	Gln
	Glu	Asn 1010		Leu	ı Let	ı Lys			nr A	la Pi	ro A			Arg	Ala	Gln
420		1010)			_	101	L5				1	020	_	_	
420			Leu			_	101	l5 J As				1 ln G	020	_	Ala Ala	
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420 423 424	Glu	1010 Asn 1025	Leu Glu	ı Lys	. Thr	· Val	101 Arg 103	L5 J As BO B G:	sp Ly	ys Pl	he G	ln G 1 le C	020 lu 035	Ser	Ala	Asp
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420 423 424 427 428 431 432 435 436 439	Glu Val Phe Phe	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg	Leu Glu Glr His	ı Lys ı Ala ı Val	S Thr Ser Lys	Val Arg Arg	101 Arg 103 Lys 104 Arg 106 Ser 107	15 As 30 G: G: 45 As 50 c I: 75 As	sp Ly lu A rg Ty	ys Pl la Ai yr A: sp G:	he G rg I sp A	ln G le C la P la P le T le T	020 lu 035 ys 050 he 065 yr 080	Ser Arg Ser Lys	Ala Gln Gln	Asp Glu Cys Leu
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420 423 424 427 428 431 432 435 436 439 440 443	Glu Val Phe Phe Cys Glu	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu	Leu Glu Glr His Asr	Lys Ala Val S Ile Asn Tyr	S Thr Ser Lys Ser Ser Lev	Val Arg Arg Val Ala	101 Arg 103 104 Arg 106 Ser 107 Glr 109	15 As	sp Ly lu Al rg Ty le As la Pl	ys Pl la Ar yr Ar sp G: ne Le	ne Girg II sp Ai ln II eu Se	ln G le C la P la P le T	020 lu 035 ys 050 he 065 yr 080 ro 095 ys	Ser Arg Ser Lys Glu Val	Ala Gln Gln Lys Asn	Asp Glu Cys Leu Pro
420 423 424 427 428 431 432 435 436 439 440 443	Glu Val Phe Phe Cys Glu Gly	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100	Glu Glr His Asr	Lys Ala Val S Ile Asn Tyr	S Thr Ser Lys Ser Ser Lev	Val Arg Arg Val Ala	101 Arg 103 104 Arg 106 Ser 107 Glr 109	15 As 30 As	sp Ly lu Al rg Ty le As la Pl	ys Pl la Ar yr Ar sp G: ne Le	ne Girg II sp Ai ln II eu Se	ln G le C la P la P le T le T sn C ler G	020 lu 035 ys 050 he 065 yr 080 ro 095 ys	Ser Arg Ser Lys Glu Val	Ala Gln Gln Lys Asn	Asp Glu Cys Leu Pro
420 423 424 427 428 431 432 435 436 439 440 443 444 447 448	Glu Val Phe Phe Cys Glu Gly	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys	Glu Glr His Asr	Lys Ala Val S Ile Asr Tyr	S Thr Ser Lys Ser Ser Leu	Val Arga Arga Val Ala Ala Asp	103 103 104 106 106 Ser 107 109 Gly 110 Met	15 As 30 As	sp Ly lu Ai rg Ty le Ai la Pl le Se	ys Pl la Ar yr As sp G: ne Le er T; sn Le	he Girg II sp AI ln II eu Se yr Ai	ln G le C la P la P le T le T sn C ler G	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 ly	Ser Arg Ser Lys Glu Val	Ala Gln Gln Lys Asn	Asp Glu Cys Leu Pro Pro Lys
420 423 424 427 428 431 432 435 436 439 440 443 444 447 448 451 452	Glu Val Phe Phe Cys Glu Gly Cys	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val	Glu Glr His Asr Arc	Lys Ala Val S Ile Asr O Tyr Phe	S Thr Ser Lys Ser Ser Leu Met	Val Arga Arga Val Ala Asp Pro	103 103 104 106 106 Ser 107 109 Gly 110 Met 112 Leu	15 As	sp Ly lu Ai rg Ty le Ai la Pl le Se sp Ai	ys Pl la Ar yr As sp Gi ne Le er Tr sn Le	he Girg II sp AI ln II eu Se yr Ai eu Se la Va	ln G le C la P le T le T sn C er G al H	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 ly 125 is	Ser Arg Ser Lys Glu Val Gly Ser	Ala Gln Gln Lys Asn Ala Glu Phe	Asp Glu Cys Leu Pro Pro Lys Arg
420 423 424 427 428 431 432 435 436 439 440 443 444 447 448 451 452 455	Glu Val Phe Phe Cys Glu Gly Cys	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val 1130 Ala	Glu Glr His Asr Arg	Lys Ala Val S Ile Asr O Tyr Phe	S Thr Ser Lys Ser Ser Leu Met	Val Arga Arga Val Ala Asp Pro	103 103 104 106 106 Ser 107 109 110 Met 112 Leu 113	15 As	sp Ly lu Ai rg Ty le Ai la Pl le Se sp Ai	ys Pl la Ar yr As sp Gi ne Le er Tr sn Le	he Girg II sp AI ln II eu Se yr Ai eu Se la Va	ln G le C la P le T er C er G al H sp A	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 ly 125 is	Ser Arg Ser Lys Glu Val Gly Ser	Ala Gln Gln Lys Asn Ala Glu Phe	Asp Glu Cys Leu Pro Pro Lys Arg
420 423 424 427 428 431 435 435 436 449 447 448 451 455 456	Glu Val Phe Phe Cys Glu Gly Cys Pro	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val 1130 Ala 1145	Glu Glr His Asr Arg	Lys Ala Val Sile Asn Tyr Phe Ala	S Thrace Service Met	Val Arga Arga Val Ala Asp Pro	103 103 104 104 106 Ser 107 109 110 Met 112 Leu 113	15 As	sp Ly lu A: rg Ty le A: la Pl le Se sp A: eu Pl sp G:	ys Pl la Ar yr As sp G ne Le er Ty sn Le ne Al	he G. rg I: sp A: ln I: eu Se eu Se eu Se la Va	ln G le C la P le T er C er G al H sp A	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 ly 125 is 140 la 155	Ser Arg Ser Lys Glu Val Gly Ser Ala	Ala Gln Gln Lys Asn Ala Glu Phe Leu	Asp Glu Cys Leu Pro Pro Lys Arg
420 423 424 427 428 431 432 435 436 439 440 447 448 451 455 456 459	Glu Val Phe Phe Cys Glu Gly Cys Pro	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val 1130 Ala 1145 Thr	Leu Glu Glr His Asr Pro Ala	Lys Ala Val Sile Asn Tyr Phe Ala	S Thrace Service Met	Val Arga Arga Val Ala Asp Pro	103 103 104 106 106 Ser 107 108 112 113 Leu 113 Val	15 As	sp Ly lu A: rg Ty le A: la Pl le Se sp A: eu Pl sp G:	ys Pl la Ar yr As sp Gi ne Le er Tr sn Le	he G. rg I: sp A: ln I: eu Se eu Se eu Se la Va	ln G le C la P la T er P er G al H sp A le L	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 ly 125 140 la 155 ys	Ser Arg Ser Lys Glu Val Gly Ser Ala	Ala Gln Gln Lys Asn Ala Glu Phe	Asp Glu Cys Leu Pro Pro Lys Arg
420 423 424 427 428 431 432 435 436 449 447 448 451 455 456 459 460	Glu Val Phe Phe Cys Glu Gly Cys Pro Asn	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val 1130 Ala 1145 Thr	Glu Glr His Asr Pro Ala Pro	Lys Ala Val Sile Asn Tyr Phe Ala Phe	S Thrace Service Metal Leue Phe	Val Arga Arga Val Ala Asp Ala Asp Ala Val	103 103 104 106 106 Ser 107 108 119 119 119 119 119 119	15 As	sp Lyrg Tyle As le Se As eu Pl	ys Pla Aryr Arsp G. sp G. sn Lear Tysn Lear A. lu Va	he G. rg I: sp A: ln I: eu Se yr A: eu Se la Va al A:	ln G le C la P la P le T le T le T le T le A le L le L	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 l25 is 140 la 155 ys	Ser Arg Ser Lys Glu Val Gly Ser Ala Glu	Ala Gln Gln Lys Asn Ala Glu Phe Leu Gln	Asp Glu Cys Leu Pro Pro Lys Arg Asp Ser
420 423 424 427 428 431 435 436 439 440 447 448 451 455 456 460 463	Glu Val Phe Phe Cys Glu Gly Cys Pro Asn	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val 1130 Ala 1145 Thr 1160 Glu	Leu Glu Glr His Asr Pro Ala Pro Asr	Lys Ala Val Sile Asn Tyr Phe Ala Phe	S Thrace Service Metal Leue Phe	Val Arga Arga Val Ala Asp Ala Asp Ala Val	103 103 104 104 106 Ser 107 109 110 111 112 113 114 116	L5 As	sp Lyrg Tyle As le Se As eu Pl	ys Pl la Ar yr As sp G ne Le er Ty sn Le ne Al	he G. rg I: sp A: ln I: eu Se yr A: eu Se la Va al A:	ln G le C la P la P le T er G la H sp A le L le L le L	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 l25 is 140 la 155 ys	Ser Arg Ser Lys Glu Val Gly Ser Ala Glu	Ala Gln Gln Lys Asn Ala Glu Phe Leu	Asp Glu Cys Leu Pro Pro Lys Arg Asp Ser
420 4224 427 428 431 4335 436 443 444 455 455 456 463 464	Glu Val Phe Phe Cys Glu Gly Cys Pro Asn Gln	1010 Asn 1025 Phe 1040 Glu 1055 Glu 1070 Arg 1085 Glu 1100 Lys 1115 Val 1130 Ala 1145 Thr	Glu Glu Glr His Asr Pro Ala Pro Asr	Lys Ala Val Asr Tyr Phe Ala Phe Ile	S Thrace Ser Level Met Level Phe	Val Arg Arg Val Ala Asp Ala Asp Ala Asp Lys	103 103 104 104 106 Ser 107 109 110 111 112 113 116 116	L5 As	sp Lyrg Tyle Asple Se eu Pl	ys Plla Anyr Asyr Assp Gine Lear Tysn Lear Tys	he Graphe	ln G le C la P la P le T er G al H sp A le L le L le L	020 lu 035 ys 050 he 065 yr 080 ro 095 ys 110 la 155 ys 170 ys 185	Ser Arg Ser Lys Glu Val Gly Ser Ala Glu Glu	Ala Gln Gln Lys Asn Ala Glu Phe Leu Gln Glu	Asp Glu Cys Leu Pro Pro Lys Arg Asp Ser Phe

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/570,047

DATE: 03/08/2006 TIME: 13:21:04

Input Set : A:\39363a.txt

Output Set: N:\CRF4\03072006\J570047.raw

ease Note:

of n and/or Xaa have been detected in the Sequence Listing. Please review the quence Listing to ensure that a corresponding explanation is presented in the <220> <223> fields of each sequence which presents at least one n or Xaa.

#:9; N Pos. 108,159,178,295,333

VERIFICATION SUMMARY

DATE: 03/08/2006

PATENT APPLICATION: US/10/570,047

TIME: 13:21:04

Input Set : A:\39363a.txt

Output Set: N:\CRF4\03072006\J570047.raw

10 M:270 C: Current Application Number differs, Replaced Current Application No 10 M:271 C: Current Filing Date differs, Replaced Current Filing Date

1267 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:60

341 Repeated in SeqNo=9